

# SEQUENCE LISTING

<110> APPLICANT: Ramakrishnan, Shyam

<120> TITLE: Regulation of Human Lipoxin A4  
Receptor-Like Protein

<130> DOCKET/FILE REFERENCE: 4974.00453

<150> PRIOR APPLICATION NUMBER: 60/189,037

<151> FILING DATE: 2000-03-14

<150> PRIOR APPLICATION NUMBER: PCT application (attorney docket LIO-005)

<151> FILING DATE: 2000-03-12

<160> NUMBER OF SEQUENCES: 5

<170> SOFTWARE: FastSEQ for Windows Version 4.0

<210> SEQ ID NO:1

<211> LENGTH: 1413

<212> TYPE: DNA

<213> ORGANISM: Homo sapiens

<400> SEQ ID NO:1

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ctccttgggc tgccagccaa tgggttgatg gcgtggctgg ccggctccca ggcccggcat    180
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ctgccagagc cgatggcaga ggcccagtc cagatggatc ctgtggccca gcctcaggtg 1080  
aaccacacac tccagccacg atcggtccc acagctcagc cacagctgaa ccctacggcc 1140  
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<210> SEQ ID NO:2

<211> LENGTH: 470

<212> TYPE: PRT

<213> ORGANISM:Homo sapiens

<400> SEQ ID NO:2

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Thr Val Phe Leu Val Ala Leu Leu Leu Gly Leu Pro Ala Asn Gly  
35 40 45  
Leu Met Ala Trp Leu Ala Gly Ser Gln Ala Arg His Gly Ala Gly Thr  
50 55 60  
Arg Leu Ala Leu Leu Leu Ser Leu Ala Leu Ser Asp Phe Leu Phe  
65 70 75 80  
Leu Ala Ala Ala Ala Phe Gln Ile Leu Glu Ile Arg His Gly Gly His  
85 90 95  
Trp Pro Leu Gly Thr Ala Ala Cys Arg Phe Tyr Tyr Phe Leu Trp Gly  
100 105 110  
Val Ser Tyr Ser Ser Gly Leu Phe Leu Leu Ala Ala Leu Ser Leu Asp  
115 120 125  
Arg Cys Leu Leu Ala Leu Cys Pro His Trp Tyr Pro Gly His Arg Pro  
130 135 140  
Val Arg Leu Pro Leu Trp Val Cys Ala Gly Val Trp Val Leu Ala Thr  
145 150 155 160  
Leu Phe Ser Val Pro Trp Leu Val Phe Pro Glu Ala Ala Val Trp Trp  
165 170 175  
Tyr Asp Leu Val Ile Cys Leu Asp Phe Trp Asp Ser Glu Glu Leu Ser  
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195                      200                      205  
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 Arg Gln Gln Gln Pro Ala Ala Cys Arg Gly Phe Ala Arg Val Ala Arg  
 225                      230                      235                      240  
 Thr Ile Leu Ser Ala Tyr Val Val Leu Arg Leu Pro Tyr Gln Leu Ala  
 245                      250                      255  
 Gln Leu Leu Tyr Leu Ala Phe Leu Trp Asp Val Tyr Ser Gly Tyr Leu  
 260                      265                      270  
 Leu Trp Glu Ala Leu Val Tyr Ser Asp Tyr Leu Ile Leu Leu Asn Ser  
 275                      280                      285  
 Cys Leu Ser Pro Phe Leu Cys Leu Met Ala Ser Ala Asp Leu Arg Thr  
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 Leu Leu Arg Ser Val Leu Ser Ser Phe Ala Ala Ala Leu Cys Glu Glu  
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 Glu Gly Pro Thr Leu Pro Glu Pro Met Ala Glu Ala Gln Ser Gln Met  
 340                      345                      350  
 Asp Pro Val Ala Gln Pro Gln Val Asn Pro Thr Leu Gln Pro Arg Ser  
 355                      360                      365  
 Asp Pro Thr Ala Gln Pro Gln Leu Asn Pro Thr Ala Gln Pro Gln Ser  
 370                      375                      380  
 Asp Pro Thr Ala Gln Pro Gln Leu Asn Leu Met Ala Gln Pro Gln Ser  
 385                      390                      395                      400  
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 405                      410                      415  
 Pro Ala Ala Ser Ser Val Pro Ser Pro Cys Asp Glu Ala Ser Pro Thr  
 420                      425                      430  
 Pro Ser Ser His Pro Thr Pro Gly Ala Leu Glu Asp Pro Ala Thr Pro  
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<210> SEQ ID NO:3

<211> LENGTH: 2300

<212> TYPE: DNA

<213> ORGANISM:Homo sapiens

<400> SEQ ID NO:3

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agtgagacct tgtttctact aaaaatttaa aaagtagtgg gtgcacacct gtagtcccag	180
ctactaggga ggctgagatg ggagggctgc tggaaaccag gaggtggaag ctgcagggac	240
tgtgccactg cactcatcct gggcaataga gcaaggccct gtctctcaa aaaaaaaaaa	300
agaaaagaaa agaaaagtct ggggtgagcc ctggcacctc ctttctacc ttcactgatt	360
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gatggtgact tgcctggaga tgcacagcac cgtctctccc atactcggtc attcacacca	540
tcattgattc accaggcacc cactccgtgt ccagcaggac tctggggacc ccaaatggac	600
actaccatgg aagctgacct ggggtgccact ggccacaggc cccgcacaga gcttgatgat	660
gaggactcct acccccaagg tggctgggac acggtcttcc tggtgccct gctgctcctt	720
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<210> SEQ ID NO:4  
 <211> LENGTH: 24  
 <212> TYPE: DNA  
 <213> ORGANISM:Homo sapiens

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<210> SEQ ID NO:5  
 <211> LENGTH: 24  
 <212> TYPE: DNA  
 <213> ORGANISM:Homo sapiens

<400> SEQ ID NO:5  
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